

SEQUENCE LISTING

SEQ ID NO:1

1 TGGCCCTCGA GGCCAAGAAT TCGGCACGAG GAGGCAGGGGA GCCAGAGGCG
51 CCAGGACCCCT CGCGTGGCGC TCCAGCACCC CAGACCGTGG CGGCGCCTCG
101 CCTTAGGGAA GAGCAAGGGAA AGAACTTTAT TTGAACCGCG AACATTTTT
151 GGTCACTGAG ATCGAGTCTC CCAGTGCTTT GGCTTCCCGC CTCTTATCG
201 TGGGTTTGAT CCCTGAGCTG CTCTCCTTTC CCGAACCTCC CGGGGTGCAG
251 CCTAGAGCCC TCCCGCGCGG CTGACTCCAG AGTAGAGGAA GGGAGGCAGC
301 CTCCGGCTGG TCCCCCGAAG CCCTCGCTGC CCCGCAGATG CGGATGGCCA
351 GCCAGTAGCG GGCGGTGGCC CCGCGTCCCG GGAGCGCACA GCAATGCAGG
401 CGCTTAACAT TACCCCGGAG CAGTTCTCTC GGCTGCTGCG GGACCACAAAC
451 CTGACGCGGG AGCAGTTCAT CGCTCTGTAC CGGCTGCGAC CGCTCGTCTA
501 CACCCCAGAG CTGCCGGAC GCGCCAAGCT GGCCCTCGTG CTCACCGCG
551 TGCTCATCTT CGCCCTGGCG CTCTTGCA ATGCTCTGGT GTTCTACGTG
601 GTGACCCGCA GCAAGGCCAT GCGCACCGTC ACCAACATCT TTATCTGCTC
651 CTTGGCGCTC AGTGACCTGC TCATCACCTT CTTCTGCATT CCCGTCACCA
701 TGCTCCAGAA CATTCCGAC AACTGGCTGG GGGGTGCTTT CATTGCAAG
751 ATGGTGCCAT TTGTCAGTC TACCGCTGTT GTGACAGAAA TCCTCACTAT
801 GACCTGCATT GCTGTGGAAA GGCACCAAGGG ACTTGTGCAT CCTTTAAAAA
851 TGAAGTGGCA ATACACCAAC CGAAGGGCTT TCACAATGCT AGGTGTGGTC
901 TGGCTGGTGG CAGTCATCGT AGGATCACCC ATGTGGCACG TGCAACAACT
951 TGAGATCAAA TATGACTTCC TATATGAAAA GGAACACATC TGCTGCTTAG
1001 AAGAGTGGAC CAGCCCTGTG CACCAGAAGA TCTACACCAC CTTCATCCTT
1051 GTCATCCTCT TCCTCCTGCC TCTTATGGTG ATGCTTATTC TGTACAGTAA
1101 AATTGGTTAT GAACTTTGGA TAAAGAAAAG AGTTGGGAT GGTCAGTGC
1151 TTCGAACTAT TCATGGAAAA GAAATGTCCA AAATAGCCAG GAAGAAGAAA
1201 CGAGCTGTCA TTATGATGGT GACAGTGGTG GCTCTTTG CTGTGTGCTG

1251 GGCACCATTG CATGTTGTCC ATATGATGAT TGAATACAGT AATTTGAAA
1301 AGGAATATGA TGATGTCACA ATCAAGATGA TTTTGCTAT CGTGCAAATT
1351 ATTGGATTTT CCAACTCCAT CTGTAATCCC ATTGTCTATG CATTATGAA
1401 TGAAAACCTTC AAAAAAAATG TTTTGTCTGC AGTTTGTAT TGCATAGTAA
1451 ATAAAACCTT CTCTCCAGCA CAAAGGCATG GAAATTCAAGG AATTACAATG
1501 ATGCGGAAGA AAGCAAAGTT TTCCCTCAGA GAGAATCCAG TGGAGGAAAC
1551 CAAAGGAGAA GCATTCAGTG ATGGCAACAT TGAAGTCAAA TTGTGTGAAC
1601 AGACAGAGGA GAAGAAAAAG CTCAAACGAC ATCTTGCTCT CTTTAGGTCT
1651 GAACTGGCTG AGAATTCTCC TTTAGACAGT GGGCATTAAAT TATAACAATA
1701 TCTTCATAAT

SEQ ID NO:2

1 MQALNITPEQ FSRLLRDHNL TREQFIALYR LRPLVYTPEL PGRAKLALVL
51 TGVLIFALAL FGNALVFYVV TRSKAMRTVT NIFICSLALS DLLITFFCIP
101 VTMLQNIISDN WLGGAFICKM VPFWQSTAVV TEILTMTCIA VERHQGLVHP
151 FKMKWQYTNR RAFTMLGVVV LVAVIVGSPM WHVQQLEIKY DFLYEKEHIC
201 CLEEWTSVPVH QKIYTTFILV ILFLLPLMVM LILYSKIGYE LWIKKRVGDG
251 SVLRTIHGKE MSKIARKKKR AVIMMVTVVA LFAVCWAPFH VVHMMIEYSN
301 FEKEYDDVTI KMIFAIQII GFSNSICNPI VYAFMNENFK KNVLSAVCYC
351 IVNKTFSPAQ RHGNSGITMM RKKAKFSLRE NPVEETKGEA FSDGNIEVKL
401 CEQTEEKKKL KRHLALFRSE LAENSPLDSG H*

SEQ ID NO:3

ATGCAGG

401 CGCTTAACAT TACCCCGGAG CAGTTCTCTC GGCTGCTGCG GGACCACAAAC
451 CTGACGCGGG AGCAGTTCAT CGCTCTGTAC CGGCTGCGAC CGCTCGTCTA
501 CACCCAGAG CTGCCGGGAC GCGCCAAGCT GGCCCTCGTG CTCACCGGCG
551 TGCTCATCTT CGCCCTGGCG CTCTTGCA ATGCTCTGGT GTTCTACGTG
601 GTGACCCGCA GCAAGGCCAT GCGCACCGTC ACCAACATCT TTATCTGCTC

651 CTTGGCGCTC AGTGACCTGC TCATCACCTT CTTCTGCATT CCCGTCACCA
701 TGCTCCAGAA CATTCCGAC AACTGGCTGG GGGGTGCTTT CATTGCAAG
751 ATGGTGCCAT TTGTCCAGTC TACCGCTGTT GTGACAGAAA TCCTCACTAT
801 GACCTGCATT GCTGTGGAAA GGCACCAGGG ACTTGTGCAT CCTTTTAAAAA
851 TGAAGTGGCA ATACACCAAC CGAAGGGCTT TCACAATGCT AGGTGTGGTC
901 TGGCTGGTGG CAGTCATCGT AGGATCACCC ATGTGGCACG TGCAACAAC
951 TGAGATCAAA TATGACTTCC TATATGAAAA GGAACACATC TGCTGCTTAG
1001 AAGAGTGGAC CAGCCCTGTG CACCAGAAGA TCTACACCAC CTTCATCCTT
1051 GTCATCCTCT TCCTCCTGCC TCTTATGGTG ATGCTTATTTC TGTACAGTAA
1101 AATTGGTTAT GAACTTTGGA TAAAGAAAAG AGTTGGGGAT GGTCAGTGC
1151 TTGAACTAT TCATGGAAAA GAAATGTCCA AAATAGCCAG GAAGAAGAAA
1201 CGAGCTGTCA TTATGATGGT GACAGTGGTG GCTCTTTG CTGTGTGCTG
1251 GGCACCATTG CATGTTGTCC ATATGATGAT TGAATACAGT AATTTGAAA
1301 AGGAATATGA TGATGTCACA ATCAAGATGA TTTTGCTAT CGTCAAATT
1351 ATTGGATTTC CCAACTCCAT CTGTAATCCC ATTGTCTATG CATTATGAA
1401 TGAAAACCTTC AAAAAAAATG TTTTGTCTGC AGTTTGTAT TGCATAGTAA
1451 ATAAACCTT CTCTCCAGCA CAAAGGCATG GAAATTCAAGG AATTACAATG
1501 ATGCGGAAGA AAGCAAAGTT TTCCCTCAGA GAGAATCCAG TGGAGGAAAC
1551 CAAAGGAGAA GCATTCAGTG ATGGCAACAT TGAAGTCAAA TTGTGTGAAC
1601 AGACAGAGGA GAAGAAAAAG CTCAAACGAC ATCTTGCTCT CTTTAGGTCT
1651 GAACTGGCTG AGAATTCTCC TTTAGACAGT GGGCATTAA